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US-09-764-876-20
; Sequence 20, Application US/09764876
; Publication No. US20030044890A1
; GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PTZ25
; CURRENT APPLICATION NUMBER: US/09/764,876
; CURRENT FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/179,065
; PRIOR FILING DATE: 2000-01-31
; PRIOR APPLICATION NUMBER: 60/180,628
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/214,886
; PRIOR FILING DATE: 2000-06-28
; PRIOR APPLICATION NUMBER: 60/217,487
; PRIOR APPLICATION NUMBER: 60/217,487
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/225,758
; PRIOR FILING DATE: 2000-08-14
; PRIOR FILING DATE: 2000-08-14
; PRIOR APPLICATION NUMBER: 60/217,496
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/217,496
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/217,496
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/225,447
; PRIOR FILING DATE: 2000-08-14
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32.9%; Score 2590.5; DB 11; Length 528; 94.3%; Pred. No. 6.1e-221; Query Match Best Local Similarity Matches 514; Conservative 2; Mismatches Indels 21; Gaps Qу Db 986 MPSGCGEQNMVLFAPIIYVLQYLEKAGLLTEEIRSRAVGFLEIGYQKELMYKHSNGSYSA 1045 Qy Db 1046 FGERDGNGNTWLTAFVTKCFGQAQKFIFIDPKNIQDALKWMAGNQLPSGCYANVGNLLHT 1105 Οy Db Qу Db Oy Db TKPSLTQKEIAKATSIVAWLAKQHNAYGGFSSTQDTVVALQALAKYATTAYMPSEEINLV 1285 Qу Db Qγ Db Qу Db Qу 1466 LSQYT 1470 Qу Db 520 TIQYS 524

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US-09-764-853-583
; Sequence 583, Application US/09764853
; Patent No. US20020090672A1
; GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PJZ06
CURRENT APPLICATION NUMBER: US/09/764,853
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper
NUMBER OF SEQ ID NOS: 939
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 583
LENGTH: 531
TYPE: PRT
ORGANISM: Homo sapiens
US-09-764-853-583
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33.0%; Score 2596.5; DB 9; Length 531; 94.3%; Pred. No. 1.8e-221; tive 2; Mismatches 8; Indels 21;
 Query Match
Best Local Similarity
     516; Conservative
                               21; Gaps
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Db
      Qy
Db
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Qy
    1044
Db
    Qу
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    Oy
Db
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    Qy
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Qу
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Db
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